Please replace the paragraph beginning at line 16 and ending at line 24 of page 7 with the following rewritten paragraph:

--sp|P00737|HPT1_HUMAN (SEQ ID NO:1)
sp|P00738|HPT2_HUMAN (SEQ ID NO:2)
sp|P50417|HPT_ATEGE (SEQ ID NO:3)
tr|Q60574|Q60574 (SEQ ID NO:4)
tr|Q61646|Q61646 (SEQ ID NO:5)
sp|Q62558|HPT_MUSSA (SEQ ID NO:6)
sp|P06866|HPT_RAT (SEQ ID NO:7)
tr|O35086|O35086 (SEQ ID NO:8)
sp|P19006|HPT_CANFA (SEQ ID NO:9)--

Please replace the paragraph beginning at line 34 of page 7 with the following rewritten paragraph:
--VLSPADKTNVKAAWGKVGAHAGEYGAEALERMFLSFPTTKTYFPHFDLSH
GSAQVKGHGKKVADALTNAVAHVDDMPNALSALSDLHAHKLRVDPVNFKL
LSHCLLVTLAAHLPAEFTPAVHASLDKFLASVSTVLTSKYR (SEQ ID NO:14)--

Please replace the paragraph beginning at line 4 of page 8 with the following rewritten paragraph:

--VHLTPEEKSAVTALWGKVNVDEVGGEALGRLLVVYPWTQRFFESFGDLST

PDAVMGNPKVKAHGKKVLGAFSDGLAHLDNLKGTFATLSELHCDKLHVDP

ENFRLLGNVLVCVLAHHFGKEFTPPVQAAYQKVVAGVANALAHKYH (SEQ ID

NO:15)--

Please replace the paragraph beginning at line 11 of page 8 with the following rewritten paragraph:

--VHLTPEEKTAVNALWGKVNVDAVGGEALGRLLVVYPWTQRFFESFGDLSS
PDAVMGNPKVKAHGKKVLGAFSDGLAHLDNLKGTFSQLSELHCDKLHVDP
ENFRLLGNVLVCVLARNFGKEFTPOMQAAYQKVVAGVANALAHKYH (SEO ID NO:16)--

Please replace the paragraph beginning at line 19 of page 8 with the following rewritten paragraph:
--GHFTEEDKATITSLWGKVNVEDAGGETLGRLLVVYPWTQRFFDSFGNLSS
ASAIMGNPKVKAHGKKVLTSLGDAIKHLDDLKGTFAQLSELHCDKLHVDP
ENFKLLGNVLVTVLAIHFGKEFTPEVQASWQKMVTAVASALSSRYH (SEO ID NO:17)--

Please replace the paragraph beginning at line 26 of page 8 with the following rewritten paragraph:

--ALSAEDRALVRALWKKLGSNVGVYTTEALERTFLAFPATKTYFSHLDLSP

GSSQVRAHGQKVADALSLAVERLDDLPHALSALSHLHACQLRVDPASFQL

LGHCLLVTLARHYPGDFSPALQASLDKFLSHVISALVSEYR (SEQ ID NO:18)--

Please replace the paragraph beginning at line 33 of page 8 with the following rewritten paragraph:
--SLTKTERTIIVSMWAKISTQADTIGTETLERLFLSHPQTKTYFPHFDLHP
GSAQLRAHGSKVVAAVGDAVKSIDDIGGALSKLSELHAYILRVDPVNFKL
LSHCLLVTLAARFPADFTAEAHAAWDKFLSVVSSVLTEKYR (SEQ ID NO:19)--

Please replace the paragraph beginning at line 1 of page 9 with the following rewritten paragraph:
--VHFTAEEKAAVTSLWSKMNVEEAGGEALGRLLVVYPWTQRFFDSFGNLSS
PSAILGNPKVKAHGKKVLTSFGDAIKNMDNLKPAFAKLSELHCDKLHVDP
ENFKLLGNVMVIILATHFGKEFTPEVQAAWQKLVSAVAIALAHKYH (SEQ ID NO:20)--

Please replace the paragraph beginning at line 9 of page 9 with the following rewritten paragraph:

--MVHLTPVEKSAVTAXWGKVNVDEVGGEALGRLLVVYPWTQRFFESFGDLS

TPDAVMGNPKVKAHGKKVLGAFSDGLAHLDNLKGTFATLSELHCDKLHVD

PENFRLLGNVLVCVLAHHFGKEFTPPVQAAYQKVVAGVANALAHKYH (SEQ ID

NO:21)--

Please replace the paragraph beginning at line 19 of page 13 with the following rewritten paragraph:

--In particular the fragments may comprise a sequence corresponding to aa 103-347 of sp|P00737 (SEQ ID NO:1) in Fig. 4 or to aa 162-406 of sp|P00738 (SEQ ID NO:2).--

Please replace the paragraph beginning at line 30 of page 21 with the following rewritten paragraph:
--tr|Q07898|Q07898 (SEQ ID NO:10)
tr|Q07901|Q07901 (SEQ ID NO:11)
tr|Q07900|Q07900 (SEQ ID NO:12)
tr|Q07899|Q07899 (SEQ ID NO:13)--

Please replace the paragraph beginning at line 30 of page 25 with the following rewritten paragraph:

--The domains are in one embodiment arranged as follows with respect to the CD163 sequence (SEQ ID NO:10):--

Please replace the paragraph beginning at line 16 of page 40 with the following rewritten paragraph:

--cDNA encoding SRCR domain 1-6 and SRCR domain 5-9 extended with Hind III and Xho I restriction sites were amplified by polymerase chain reactions (PCR) using fulllength HbSR cDNA (Kristiansen, M., Graversen, J.H., Jacobsen, C., Sonne, O., Hoffman, H., Law, A.S.K., and K., M.S.K. (2001) Identification of the hemoglobin scavenger receptor, Nature 409, 198-201) as template. The PCR products were subcloned into the expression vector pSecTag2B (Invitrogen, Groningen, The Netherlands) by use of the restriction sites HindIII and XhoI. Plasmids were transformed into E. coli DH5lpha cells (Clontech, Palo Alto, CA, USA), and plasmid DNA isolated and sequenced prior to transfection. The following primers were used for construction of the fragments: SRCR domain 1-6: forward 5'-caagcttggaacagacaaggagctg-3' (SEQ ID NO:22) and reverse 5'-cctcgagtcctgagcagattacagag-3' (SEQ ID NO:23). SRCR domain 5-9: forward 5'-caagcttcacagggaacccagactg-3' (SEQ ID NO:24) and reverse 5'-cctcgagatctgtgcaattcactgc-3' (SEQ ID NO:25).--

IN THE SEQUENCE LISTING

Please enter the attached Sequence Listing, numbered as pages 1-44.